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FILE 'MEDLINE, AGRICOLA, CABA, CAPLUS, BIOSIS, DISSABS, EMBASE' ENTERED  
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E BEGOVICH/AU 25

L1

9 E5

E BEGOVICH/AU 25

L2

365 E4 OR E6 OR E7 OR E8 OR E9 OR E10

L3

167 DUP REM L2 (198 DUPLICATES REMOVED)

L4

25 L3 AND (RA OR ARTHRITIS)

L5

444 (PTPN22 OR HCV16021387 OR RS2476601)

L6

200 L5 AND ARTHRITIS

L7

90 DUP REM L6 (110 DUPLICATES REMOVED)

L8

77 L7 AND (ASSOCIATED OR ASSOCIATE OR ASSOCIATION)

L9

0 L8 AND PY<2004

L10

2 L8 AND PY<2005

L11

0 L7 AND PT<2004

L12

0 L7 AND PY<2004

L13

35 RS2476601

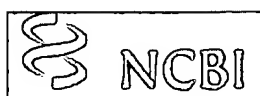
L14

20 L13 AND RA

L15

5 DUP REM L14 (15 DUPLICATES REMOVED)

12-5-06



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☐ 1: Rheumatology (Oxford). 2006 Nov 29; [Epub ahead of print]



Full Text  
Rheumatology

Links

**The 620W allele is the PTPN22 genetic variant conferring susceptibility to RA in a Dutch population.**

**Wesoly J, Hu X, Thabet MM, Chang M, Uh H, Allaart CF, Toes RE, Houwing-Duistermaat JJ, Begovich AB, Huizinga TW.**

Department of Rheumatology, Leiden University Medical Center, Leiden, The Netherlands, Celera Diagnostics, Alameda, CA, USA and Department of Medical Statistics, Leiden University Medical Center, Leiden, The Netherlands.

**OBJECTIVES:** A missense SNP, C1858T, in PTPN22 has been identified as a genetic risk factor for rheumatoid arthritis (RA). Subsequent work has suggested that other variants in this gene, in particular a haplotype marked by the minor allele of rs3789604, are associated with RA in white North Americans independent of C1858T. We tested this hypothesis in an independent white Dutch study. **METHODS:** A total of 667 RA patients and 286 controls were genotyped for 13 PTPN22 single nucleotide polymorphisms (SNPs) by allele-specific kinetic polymerase chain reaction. rs3789604 was genotyped in an additional 410 RA and 270 UA patients participating in the Leiden early arthritis inception cohort. We conducted single-marker and haplotype association tests. **RESULTS:** The sole haplotype strongly associated with RA in our Dutch population carries the PTPN22 1858T allele. A second haplotype identical at all other SNPs tested except 1858 was not associated with disease. No significant association of the haplotype tagged by the 3' PTPN22 SNP, rs3789604, with RA susceptibility ( $P = 0.134$ ) was observed in our sample set. **CONCLUSION:** We conclude that C1858T is the sole PTPN22 variant predisposing to RA in our white Dutch sample set.

PMID: 17135225 [PubMed - as supplied by publisher]

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PTPN22 genetic variation: evidence for multiple variants associated with rheumatoid arthritis. [Am J Hum Genet. 2005]

Association of the lymphoid tyrosine phosphatase R620W variant with rheumatoid arthritis, but not Crohn's disease, in Canadian patients. [Arthritis Rheum. 2005]

Rheumatoid arthritis association with the FCRL3 -169C polymorphism is restricted to PTPN22 1858T-homozygous individuals in a Canadian population. [Arthritis Rheum. 2006]

Association of PTPN22 1858 single-nucleotide polymorphism with rheumatoid arthritis in a German cohort: higher frequency of the risk allele in male compared to female patients. [Res Ther. 2006]

Haplotype analysis revealed no association between the PTPN22 gene and RA in a Japanese population. [Rheumatology (Oxford). 2006]

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L11	50	begovich.in.	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT	OR	ON	2006/12/05 09:41
L12	18	begovich.in. and ann.in.	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT	OR	ON	2006/12/05 10:04
L13	9	PTPN22 OR HCV16021387 OR RS2476601	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT	OR	ON	2006/12/05 10:02
L14	9	PTPN22 OR *16021387 OR *2476601	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT	OR	ON	2006/12/05 09:58
L15	0	*2476601	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT	OR	ON	2006/12/05 10:02
L16	0	??2476601	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT	OR	ON	2006/12/05 10:02
L17	10	"2476601"	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT	OR	ON	2006/12/05 10:02
L18	0	"16021387"	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT	OR	ON	2006/12/05 10:02
L19	0	"*1387"	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT	OR	ON	2006/12/05 10:03

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